

SEQUENCE LISTING

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Guiltinan, Mark
Shcherban, Tatyana
Jun, Shi

<120> Purified Plant Expansin Proteins and DNA Encoding Same

<130> P04666US08

<140> US

<141> 2003-09-24

<150> US 09/896,301

<151> 2001-06-29

<150> US 09/092,160

<151> 1998-06-05

<150> US 09/429,675

<151> 1999-10-28

<150> US 08/834,327

<151> 1997-04-15

<150> US 08/444,515

<151> 1995-05-12

<150> US 08/242,090

<151> 1994-05-12

<150> US 08/060,944

<151> 1993-05-12

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gtggcgctga gcactgcgct atttaacaat ggattaagtt gtggtgcttg cttcgaaatg 180

acttgtaaa acgaccctaa atggtgcctt cggggaacta ttaggggtcac tgccaccaac 240

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aatggccact catacttcaa cctcgttttg atcacaaaacg tcggtggcgc aggcgacgtc 480
cactctgtgt cgataaaggg gtctcgaact ggatggcaat ccatgtctag aaattggggc 540
caaaactggc aaagcaacaa ctatctcaat ggccaaggcc tttcctttca agtcactctt 600
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Asp Ala Ser Gly Thr Met Gly Gly Ala Cys Gly Tyr Gly Asn Leu Tyr
20           25           30

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Ser Gln Gly Tyr Gly Thr Asn Thr Ala Ala Leu Ser Thr Ala Leu Phe
35           40           45

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Asn Asn Gly Leu Ser Cys Gly Ala Cys Phe Glu Ile Arg Cys Gln Asn
50           55           60

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Asp Gly Lys Trp Cys Leu Pro Gly Ser Ile Val Val Thr Ala Thr Asn
65           70           75           80

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Phe Cys Pro Pro Asn Asn Ala Leu Pro Asn Asn Ala Gly Gly Trp Cys
85           90           95

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Asn Pro Pro Gln Gln His Phe Asp Leu Ser Gln Pro Val Phe Gln Arg
100          105          110

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Ile Ala Gln Tyr Arg Ala Gly Ile Val Pro Val Ala Tyr Arg Arg Val
 115 120 125

Pro Cys Val Arg Arg Gly Gly Ile Arg Phe Thr Ile Asn Gly His Ser
 130 135 140

Tyr Phe Asn Leu Val Leu Ile Thr Asn Val Gly Gly Ala Gly Asp Val
 145 150 155 160

His Ser Ala Met Val Lys Gly Ser Arg Thr Gly Trp Gln Ala Met Ser
 165 170 175

Arg Asn Trp Gly Gln Asn Trp Gln Ser Asn Ser Tyr Leu Asn Gly Gln
 180 185 190

Ser Leu Ser Phe Lys Val Thr Thr Ser Asp Gly Gln Thr Ile Val Ser
 195 200 205

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Ala His Val Arg
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Ser Gln Gly Tyr Gly Leu Glu Thr Ala Ala Leu Ser Thr Ala Leu Phe
35 40 45

Asp Gln Gly Leu Ser Cys Gly Ala Cys Xaa Glu Leu Met Cys Val Asn
50 55 60

Asp Pro Gln Trp Cys Ile Ala Gly Arg Ser Ile Val Val Thr Ala Thr
65 70 75 80

Asn Phe Cys Pro Pro Gly Gly Ala Cys Asp Pro Pro Asn His His Phe
85 90 95

Asp Leu Ser Gln Pro Ile Tyr Glu Lys Ile Ala Leu Tyr Lys Ser Gly
100 105 110

Ile Ile Pro Val Met Tyr Arg Arg Val Arg Cys Lys Arg Ser Gly Gly
115 120 125

Ile Arg Phe Thr Ile Asn Gly His Ser Tyr Phe Asn Leu Val Leu Val
130 135 140

Thr Asn Val Gly Gly Ala Gly Asp Val His Ser Val Ser Met Lys Gly
145 150 155 160

Ser Arg Thr Lys Trp Gln Leu Met Ser Arg Asn Trp Gly Gln Asn Trp
165 170 175

Gln Ser Asn Ser Tyr Leu Asn Gly Gln Ser Leu Ser Phe Val Val Thr
180 185 190

Thr Ser Asp Arg Arg Ser Val Val Ser Phe Asn Val Ala Pro Pro Thr
195 200 205

Trp Ser Phe Gly Gln Thr Tyr Thr Gly Gly Gln Phe Arg Tyr

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20 25 30		
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35	40	45
Asn Asp Gly Ala Met Cys Gly Ala Cys Tyr Thr Ile Thr Cys Asp Thr		
50	55	60
Ser Gln Thr Lys Trp Cys Lys Pro Gly Gly Asn Ser Ile Thr Ile Thr		
65	70	75
Ala Thr Asn Leu Cys Xaa Pro Asn Trp Ala Leu Pro Ser Asn Ser Gly		
85	90	95
Gly Trp Cys Asn Pro Pro Leu Xaa His Phe Asp Met Ser Gln Pro Ala		
100	105	110
Trp Glu Asn Ile Ala Val Tyr Gln Ala Gly Ile Val Pro Val Asn Tyr		
115	120	125
Lys Arg Val Pro Xaa Gln Arg Ser Gly Gly Ile Arg Phe Ala Ile Ser		
130	135	140
Gly His Asp Tyr Phe Glu Leu Val Thr Val Thr Asn Val Gly Gly Ser		
145	150	155
Gly Val Val Ala Gln Met Ser Ile Lys Gly Ser Asn Thr Gly Trp Met		
165	170	175
Ala Met Ser Arg Asn Trp Gly Ala Asn Trp Gln Ser Asn Ala Tyr Leu		
180	185	190
Ala Gly Gln Ser Leu Ser Phe Ile Val Gln Leu Asp Asp Gly Arg Lys		
195	200	205
Val Thr Ala Trp Asn Xaa Ala Pro Xaa Asn Trp Leu Xaa Xaa Xaa Xaa		
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20 25 30

Ser Gln Gly Tyr Gly Leu Gln Thr Ala Ala Leu Ser Thr Ala Leu Phe
35 40 45

Asn Ser Gly Gln Lys Cys Gly Ala Cys Phe Glu Leu Thr Cys Glu Asp
50 55 60

Asp Pro Glu Trp Cys Ile Pro Gly Ser Ile Ile Val Arg Thr Asn Leu
65 70 75 80

Ala Asn Phe Ala Leu Ala Asn Asp Asn Gly Gly Trp Cys Asn Pro Pro
85 90 95

Leu Lys His Phe Asp Leu Ala Glu Pro Ala Phe Leu Gln Ile Ala Gln
100 105 110

Tyr Arg Ala Gly Ile Val Pro Val Ala Phe Arg Arg Val Pro Cys Glu
115 120 125

Lys Gly Gly Gly Ile Arg Phe Thr Ile Asn Gly Asn Pro Tyr Phe Asp
130 135 140

Leu Val Leu Ile Thr Asn Val Gly Gly Ala Gly Asp Ile Arg Ala Val
145 150 155 160

Ser Leu Lys Gly Ser Lys Thr Asp Gln Trp Gln Ser Met Ser Arg Asn
165 170 175

Trp Gly Gln Asn Trp Gln Ser Asn Thr Tyr Leu Arg Gly Gln Ser Leu
180 185 190

Ser Phe Gln Val Thr Asp Ser Asp Gly Arg Thr Val Val Ser Tyr Asp
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Val Val Pro His Asp Trp Gln Phe Gly Gln Thr Phe Glu Gly Gly Gln
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Phe
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Ser Thr Gly Tyr Thr Asn Thr Ala Ala Leu Ser Thr Val Leu Phe Asn
 35 40 45

Asp Gly Ala Ala Cys Arg Ser Cys Tyr Glu Leu Arg Cys Asp Asn Asp
 50 55 60

Gly Gln Trp Cys Leu Pro Gly Ser Val Thr Val Thr Ala Thr Asn Leu
 65 70 75 80

Cys Pro Pro Asn Tyr Ala Leu Pro Asn Asp Asp Gly Gly Trp Cys Asn
 85 90 95

Pro Pro Arg Pro His Phe Asp Met Ala Glu Pro Ala Phe Leu Gln Ile
 100 105 110

Gly Val Tyr Arg Ala Gly Ile Val Pro Val Ser Tyr Arg Arg Val Pro
 115 120 125

Cys Val Lys Lys Gly Gly Ile Arg Phe Thr Ile Asn Gly His Ser Thr
 130 135 140

Phe Asn Leu Val Leu Val Thr Asn Val Ala Gly Pro Gly Asp Val Gln
 145 150 155 160

Ser Val Ser Ile Lys Gly Ser Ser Thr Gly Trp Gln Pro Met Ser Arg
165 170 175

Asn Trp Gly Gln Asn Trp Gln Ser Asn Ser Tyr Leu Asp Gly Gln Ser
180 185 190

Leu Ser Phe Gln Val Ala Val Ser Asp Gly Arg Thr Val Thr Ser Asn
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Asn Val Val Pro Ala Gly Trp Gln Phe Gly Gln Thr Phe Glu Gly Gly
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Gln Phe
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Ser Gln Gly Tyr Gly Thr Asn Thr Val Ala Leu Ser Thr Ala Leu Phe
35 40 45

Asn Asn Gly Leu Ser Cys Gly Ala Cys Phe Glu Met Thr Cys Thr Asn
50 55 60

Asp Pro Lys Trp Cys Leu Pro Gly Thr Ile Arg Val Thr Ala Thr Asn
65 70 75 80

Phe Cys Pro Pro Asn Phe Ala Leu Pro Asn Asp Asp Gly Gly Trp Cys
85 90 95

Asn Pro Pro Leu Gln His Phe Asp Met Ala Glu Pro Ala Phe Leu Gln
100 105 110

Ile Ala Gln Tyr Arg Ala Gly Ile Val Pro Val Ser Phe Arg Arg Val
 115 120 125

Pro Cys Met Lys Lys Gly Gly Val Arg Phe Thr Ile Asn Gly His Ser
 130 135 140

Tyr Phe Asn Leu Val Leu Ile Thr Asn Val Gly Gly Ala Gly Asp Val
 145 150 155 160

His Ser Val Ser Ile Lys Gly Ser Arg Thr Gly Trp Gln Ser Met Ser
 165 170 175

Arg Asn Trp Gly Gln Asn Trp Gln Ser Asn Asn Tyr Leu Asn Gly Gln
 180 185 190

Gly Leu Ser Phe Gln Val Thr Leu Ser Asp Gly Arg Thr Leu Thr Ala
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Tyr Asn Leu Val Pro Ser Asn Trp Gln Phe Gly Gln Thr Tyr Glu Gly
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Pro Gln Phe
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Ala

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Cys

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Gly

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Gly Gly Gly Cys
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Cys Cys Asn Cys Cys Asn Cys Lys Arg Thr Ala Asp Ala Thr Asn Cys
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